

FIGURE 1

```
TPVgp38aa    1                               TLKYCYTVTLKDNGLYDKVFYCHYN    25
Yabagp38     1 MNKLILSLG FVATCNCITLRYNYTVTVK-NGLYDGVFFDYNDQLVTRI 49
               **.* ****.* ***** **.* **

TPVgp38aa    26                               25 (SEQ ID NO: 1)
Yabagp38     50 SYNHETKRG NVN 61 (SEQ ID NO: 2)
```

TPVgp38aa 1 TLKYCYTVTLKDNGLYDKVFYCHYN 25

FIGURE 2

YMTV partial gp38 gene (183 nucleotide):

5'
ATGAATAAGTTAATTTTATCGTTGTTGGGTTTTGTGGCAACTTGCAATTGTATAACCTTAAGATATAATTATACCGTTA
CGGTAAAGAATGGATTATACGACGGGGTATTTTTTGATTATTACAACGATCAGTTAGTAACGAGGATATCATATAATCA
TGAAACCAAACGAGGAAATGTAAAT (SEQ ID NO: 3)

YMTV partial gp38 gene (61 amino acid):

5'
MNKLILSLLGFVATCNCITLRYNYTVTVKNGLYDGVFFDYNDQLVTRISYNHETKRGNVN (SEQ ID NO: 2)

SEQ ID NO: 4

MNKLILFSTIVAVCNCITLKYNITVTLKDNGLYDGVFYDHYNDQLVTKISYNHETRHGNNVFRADWFKIS
RSPHTPGNDYNFNFWYSLMKETLEEINKNDSTKTTSLSLITGCIYETGLLFGSYGYVETANGPLARYHTGD
KRFTKMTHKGF PKVGMLTVKNTLWKDVKTYLGGFEYMGCSLAILDYQKMAKGEIPKDTTPTVKVTGNELE
DGNMTLECSVNSFYPPDVITKWIESEHFKGEYKYVNGRYYPEWGRKSDYEPGEPGF PWNKKDKDANTYS
LTDLVRTTSKMSSQLVCVVFHDTLEAQVYTCSEGCNGELYDHLRYKTEEGEGEEDEED*

FIG. 3

1000 900 800 700 600 500 400 300 200 100 0

SEQ ID NO: 5

Tana gp38:

AAGCTTCATGAATAAGTTAATATTATTTAGCACAAATTGTAGCAGTTTGTA
ACTGCATAACTTTAAAATATAATTATACTGTTACGTTAAAAGATAATGGGTTATAC
GATGGAGTATTTTACGATCATTACAACGATCAGTTAGTAACGAAAATATCAT
ATAACCACGAACTAGACACGGAAACGTAAATTTTAGGGCTGATTGGTTTAA
TATTTCTAGGAGTCCCCACACGCCAGGTAACGATTACAACTTTAACTTTTGGT
ATTCTTTAATGAAAGAACTTTAGAAGAAATTAATAAAAAACGATAGCACAAA
AACTACTTCGCTTTCATTAATCACTGGGTGTTATGAAACAGGATTATTATTG
GTAGTTATGGGTATGTAGAAACGGCCAACGGACCGTTGGCCAGATACCATAC
AGGAGATAAAAGGTTTACGAAAATGACACATAAAGGTTTTCCCAAGGTTGGA
ATGTTAACTGTAAAAAACACTCTTTGGAAAGATGTAAAAACTTATCTAGGCG
GTTTTGAATACATGGGATGTTTATTAGCTATTTTAGATTACCAAAAAATGGCT
AAAGGTGAAATACCAAAAGATACAACACCTACAGTGAAAGTAACGGGTAAAT
GAGTTAGAAGATGGTAACATGACTCTTGAATGCAGTGTAAATTCATTTTACCC
TCCTGACGTAATTACTAAGTGGATAGAAAGCGAACATTTTAAAGGTGAATAT
AAATATGTTAACGGAAGATACTATCCAGAATGGGGGAGAAAATCCGATTATG
AGCCAGGAGAGCCAGGTTTTCCATGGAATATTAATAAAAGATAAAGATGCAA
ACACATATAGTTTAACAGATTTAGTACGTACAACATCAAAAATGAGTAGTCA
ACTAGTATGTGTTGTTTTCCATGACACTTTAGAAGCGCAAGTTTATACTTGTT
CTGAAGGATGCAATGGAGAGCTATACGACCACCTATATAGAAAAACAGA
AGAAGGAGAAGGTGAAGAGGATGAAGAAGACGGAAACCCTCGAG

FIG. 4

Variable	Mean	SD	Min	Max	Skewness	Kurtosis	Normality
Age	38.5	12.5	25	65	0.1	3.2	0.95
Gender	1.2	0.4	1	2	0.0	3.0	0.98
Education	12.5	2.5	9	16	0.2	3.5	0.92
Income	1500	500	1000	2500	0.3	3.8	0.88
Health	2.5	0.5	1	3	0.1	3.1	0.96
Stress	3.5	1.0	1	5	0.4	4.0	0.85
Depression	2.0	0.8	1	4	0.2	3.3	0.93
Life Satisfaction	4.0	1.0	1	5	-0.1	3.0	0.97
Resilience	3.0	0.8	1	4	0.1	3.2	0.94
Optimism	3.5	0.9	1	4	0.0	3.1	0.96
Self-Esteem	3.0	0.7	1	4	0.1	3.2	0.95
Emotional Stability	3.5	0.8	1	4	0.0	3.1	0.97
Life Satisfaction	4.0	1.0	1	5	-0.1	3.0	0.97
Resilience	3.0	0.8	1	4	0.1	3.2	0.94
Optimism	3.5	0.9	1	4	0.0	3.1	0.96
Self-Esteem	3.0	0.7	1	4	0.1	3.2	0.95
Emotional Stability	3.5	0.8	1	4	0.0	3.1	0.97

FIG. 5

FIG. 5

SEQ ID NO: 7

YLD gp38:

ATGGATAAGTTACTATTATTTAGCACAATTGTAGCAGTTTGTAAGTGCATAAC
TTTAAAATATAATTATACTGTTACGTTAAAAGATGATGGGTTATACGATGGAG
TATTTTACGATCATTACAACGATCAGTTAGTGACGAAAATATCATATAACCAT
GAACTAGACACGGAAACGTAAATTTAGGGCTGATTGGTTTAATATTTCTA
GGAGTCCCCACACGCCAGGTAACGATTATAACTTTAACTTTTGGTATTCTTTA
ATGAAAGAACTTTAGAAGAAATTAATAAAAACGATAGCACAAAACTACTT
CGCTTTCATTAATCACTGGGTGTTATGAAACAGGATTATTATTTGGTAGTTAT
GGGTATGTAGAAACGGCCAACGGGCCGTTGGCCAGATACCATACAGGAGAT
AAAAGGTTTACGAAAATGACACATAAAGGTTTTCCCAAGGTTGGAATGTTAA
CTGTAAAAAACACTCTTTGGAAAGATGTAAAAGCTTATTTAGGCGGTTTTGA
ATATATGGGATGTTTCATTAGCTATTTTAGATTACCAAAAAATGGCTAAAGGTA
AAATACCAAAAGATACAACACCTACAGTGAAAGTAACGGGTAATGAGTTAG
AAGATGGTAACATGACTCTTGAATGCACTGTAAATTCATTTTACCCTCCTGAC
GTAATTACTAAGTGGATAGAAAGCGAACATTTTAAAGGTGAATATAAATATG
TTAACGGAAGATACTATCCAGAATGGGGGAGAAAATCCAATTATGAGCCAGG
AGAGCCAGGTTTTCCATGGAATATCAAAAAAGATAAAGATGCAAATACATAT
AGTTTAACAGATTTAGTACGTACAACATCAAAAATGAGTAGTCAACCAGTAT
GTGTTGTTTTCCATGACACTTTAGAAGCGCAAGTTTATACTTGTTCTGAAGGA
TGCAATGGAGAGCTATACGATCACCTATATAGAAAAACAGAAGAAGGG
GAAGGTGAAGAGGATGAAGAAGACTGA

FIG. 6

SEQ ID NO: 8

MITKAIVILSIITAYVDASAFLVYNYTYTLQDDNHRYDFEVTDYFNDILIKRLKLNSETGRPELRNEPPT
WFNETKIRYYPKNNYNFMFWLNRMSETLDEINKLPETSNPYKTMSLTIGCTDLRQLQVNFQYVTVGGNIW
TRFDPKNKRFSKVRSRTPFKVGMLTVKSQHWERVMHLGSMVTLTCPFTADDYYKISKGYIDKPVKPTVT
VTGIERGDNTTLICTFDNHYPSSVAVKWYNIEDFAPDYRYDPYVNELLPDTDYLPGEFGYPTITRRLGDK
YLFTSSPRVMVPTIMSNRIACVGFHSTLEPSIYRCVNCSGPEPVLQYQGDRRNDLEDEED

FIG. 7

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SEQ ID NO: 9

Swinepox C1L

ATGATTACTAAAGCGATTGTGATATTGTCTATTATTACAGCATATGTAGATGC
TTCCGCATTCTTAGTATACAATTATACATATACTTTACAAGATGATAATCATC
GATATGACTTCGAAGTCACCGATTATTTTAATGATATACTAATAAAACGTTTA
AAACTAAATAGCGAGACAGGAAGACCAGAATTAAGAAATGAACCACCAACA
TGGTTTAATGAGACTAAGATTAGATATTATCCGAAAAATAATTATAATTTTAT
GTTCTGGCTAAATAGAATGAGTGAAACGCTAGATGAGATAAAATAAACTTCCA
GAAACGAGTAATCCTTACAAGACTATGTCCTTGACAATTGGATGTACTGATCT
AAGACAACCTCAAGTAAATTTTCGGTTATGTTACTGTAGGTGGTAATATATGGA
CACGATTCGACCCCAAGAATAAACGCTTTAGTAAAGTTAGATCACGTACATT
TCCAAAGGTAGGAATGTAACTGTAAATCACAACACTGGGAACGTGTTATG
GAACATCTTGGATCAATGGTAACATTAACATGTCCGTTTACAGCGGATGATTA
TTATAAAATTTCTAAGGGATATATAGATAAGCCAGTTAAGCCTACTGTTACAG
TTACAGGAATTGAAAGAGGAGATAATACTACATTGATATGCACATTTGATAA
TCATTATCCGTCGTCGGTCGCTGTAAATGGTATAACATCGAGGACTTTGCTC
CGGACTATCGTTATGATCCGTACGTAAATGAATTGCTTCCTGATACGGACTAT
CTACCGGGTGAACCAGGATATCCGACTATAACTAGGAGATTAGGTGATAAAT
ATTTATTTACATCATCACCTAGGGTTATGGTACCAACTATCATGTCTAATAGA
ATAGCATGTGTTGGATTTCATAGTACGTTAGAACCAAGCATATATAGATGTGT
AAACTGCTCGGGACCTGAGCCTGTTTTACAATACCAGGGAGAT
AGAAGGAATGACTTGGAGGATGAGGAGGATTAA

FIG. 8

ClustalW Formatted Alignments

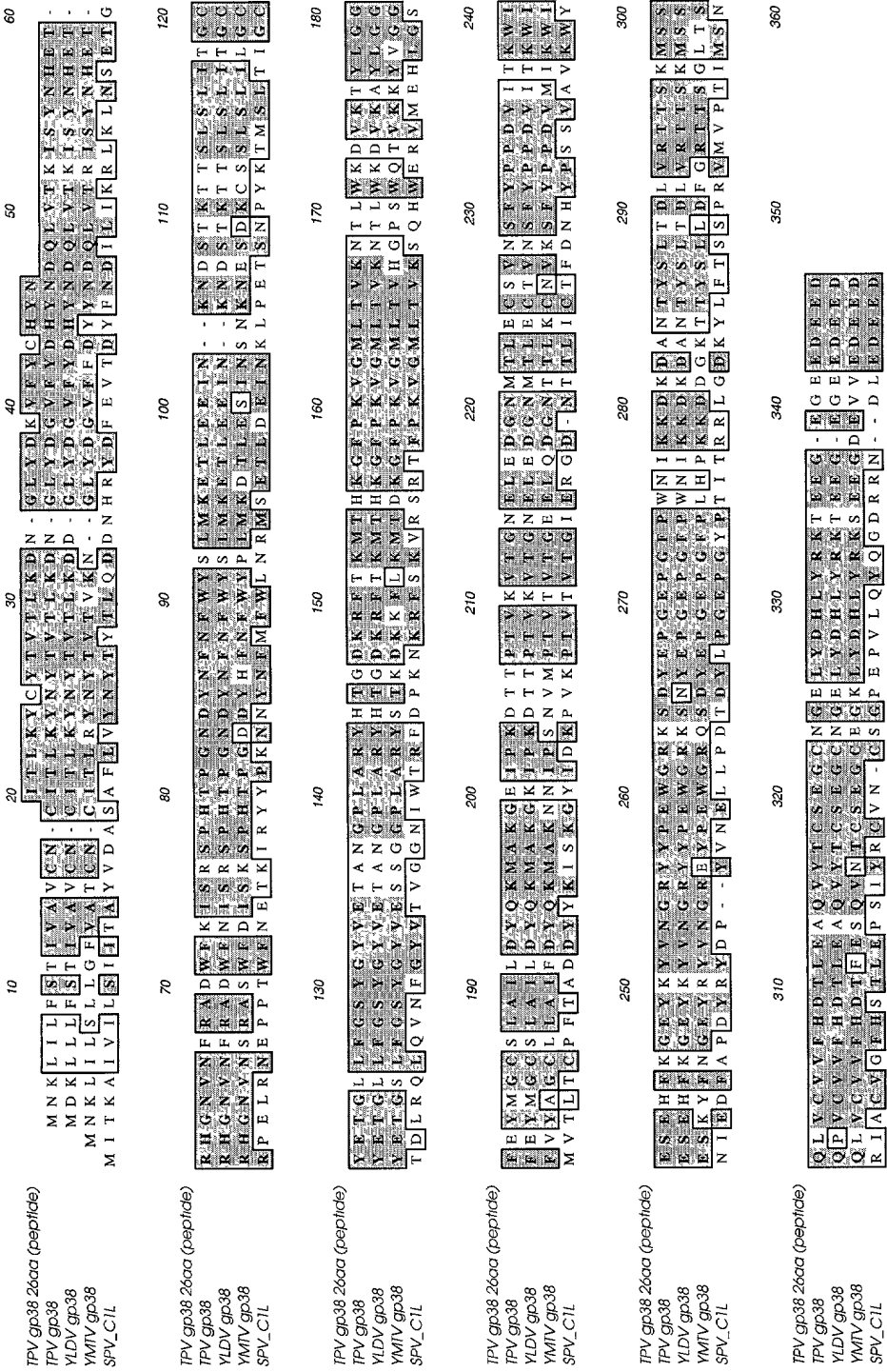


FIG. 9